

090245-08301  
FOETED-09482660

Site for insertion of DNA segment

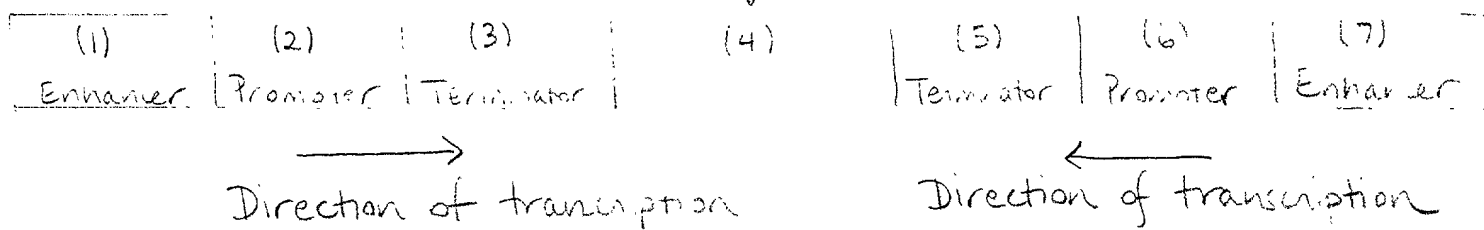


Figure 1

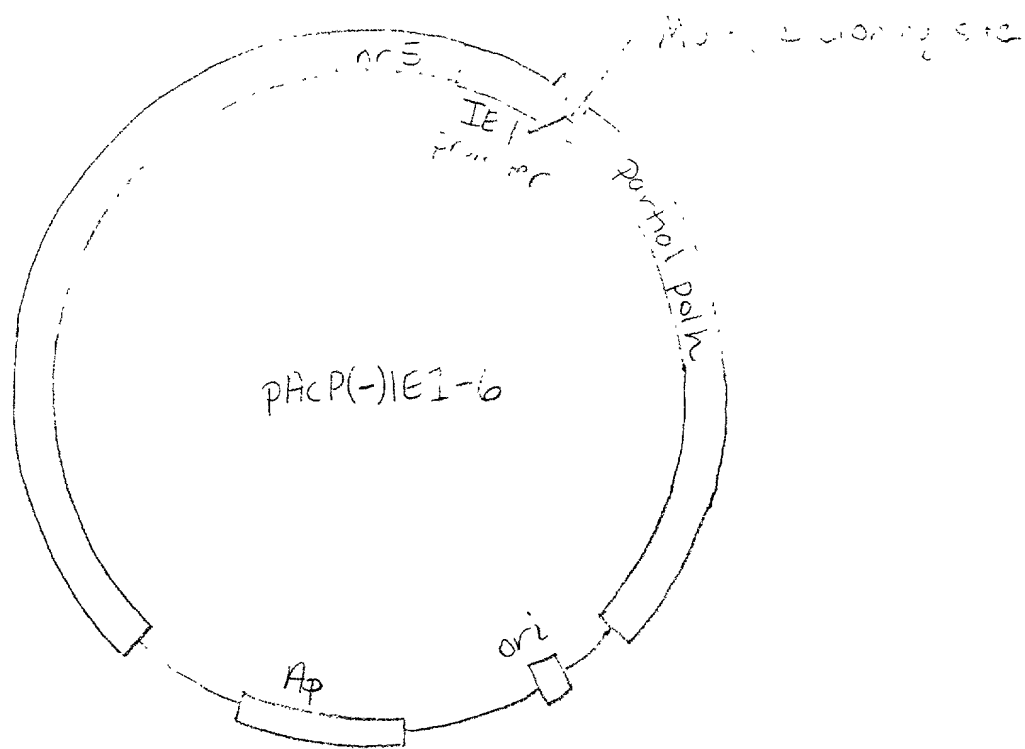
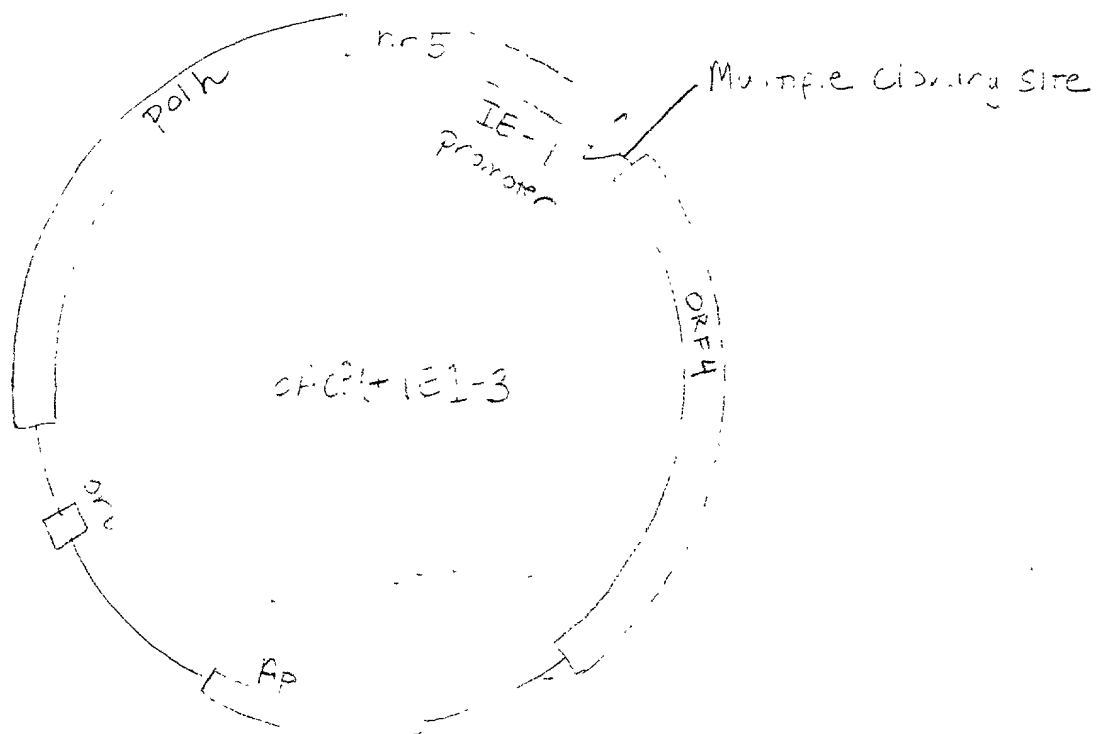


Figure 2

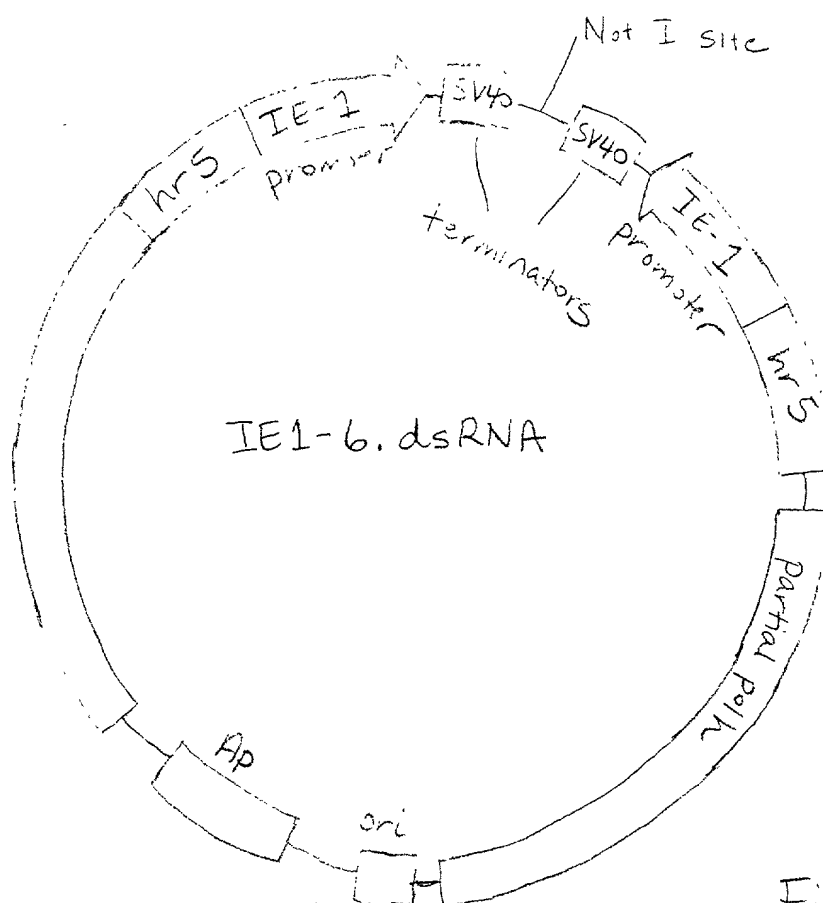
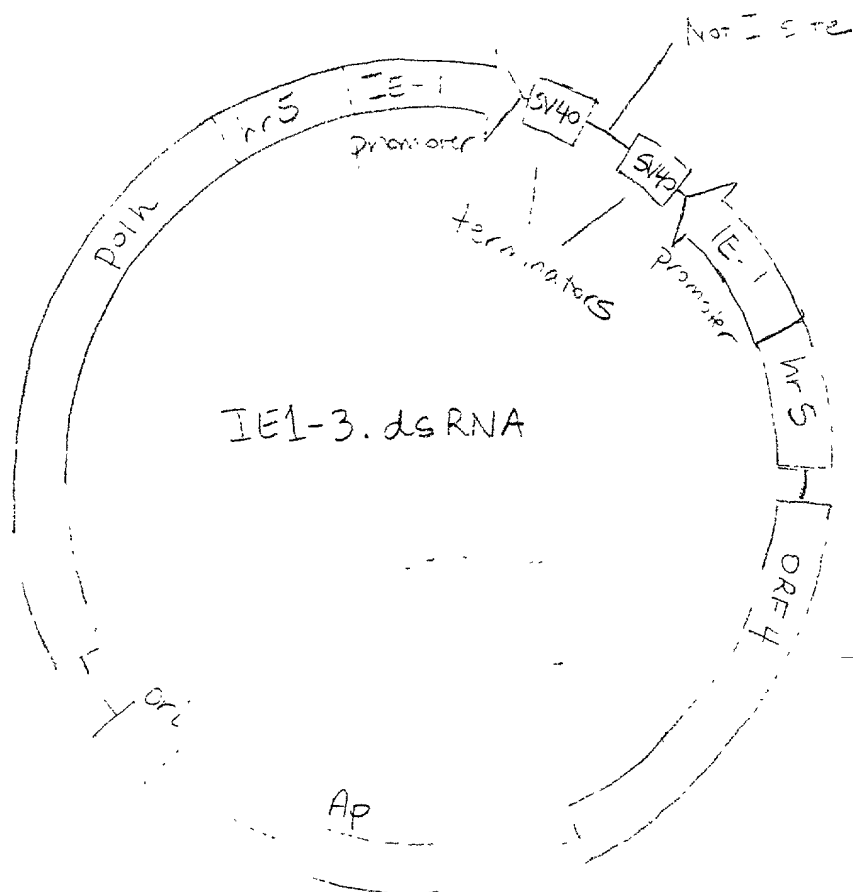


Figure 3

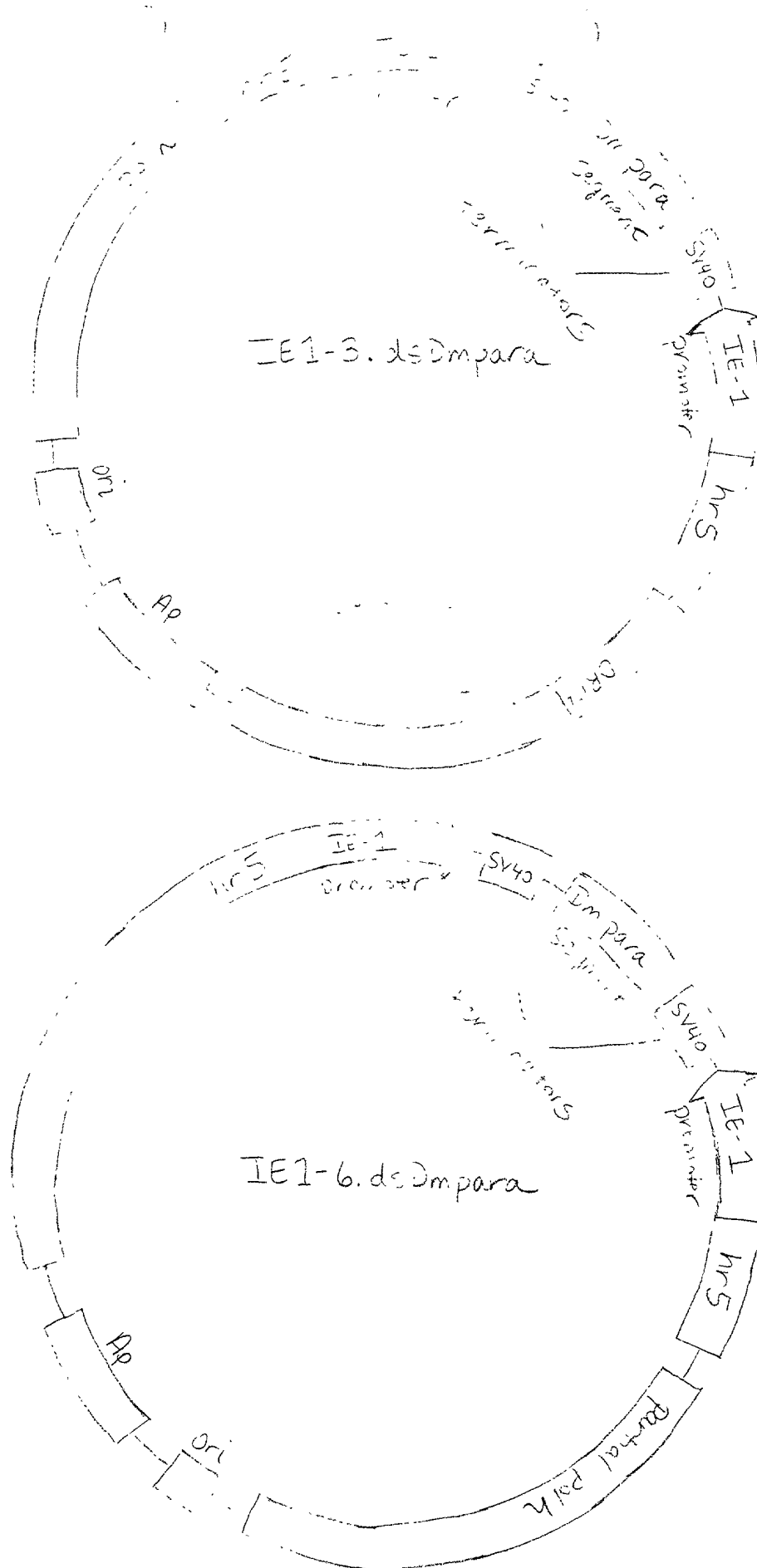


Figure 4



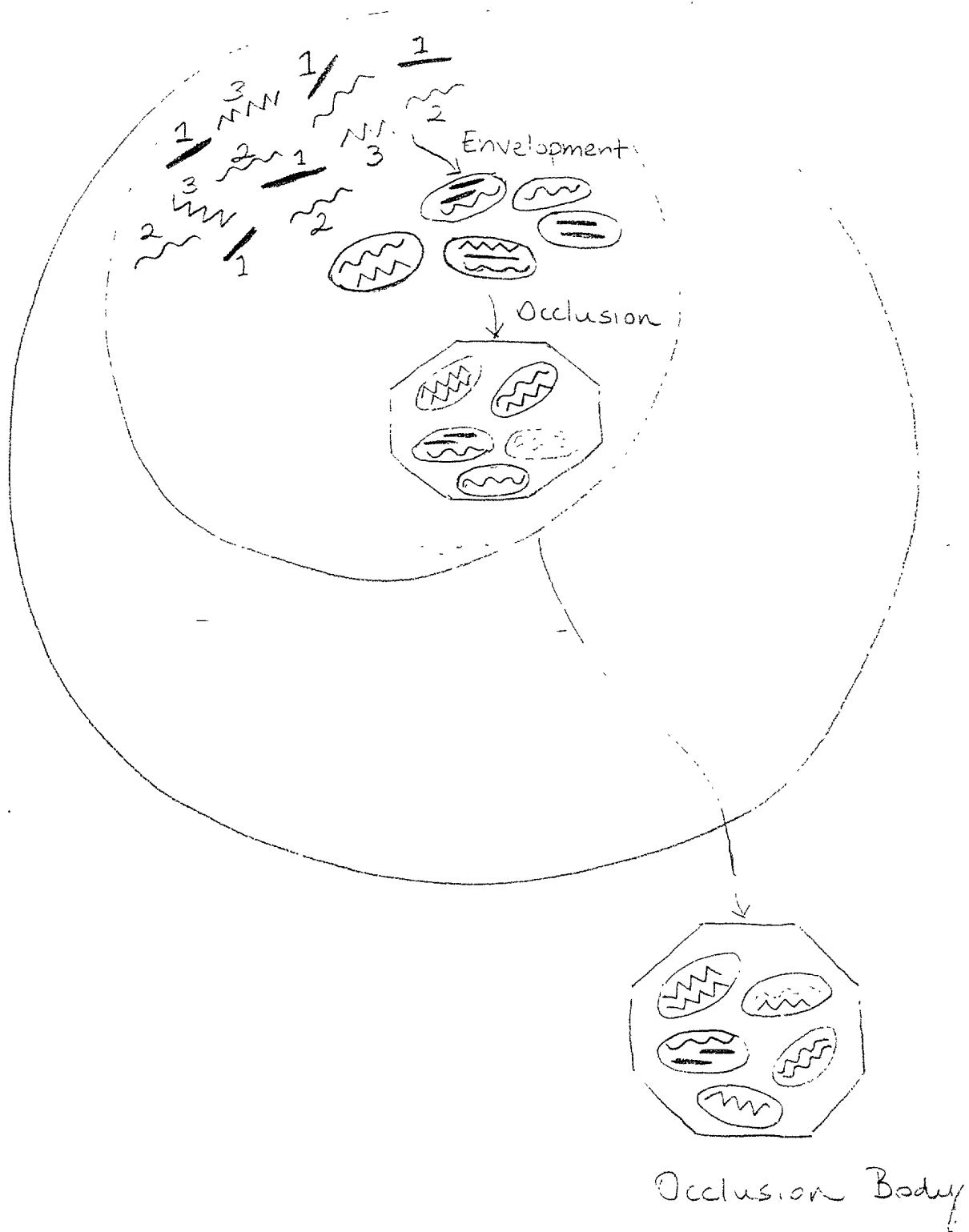


Figure 6

## Figure 7

### SV40 Transcriptional Terminator Sequence

SEQ ID NO:1

TGATCATAATCAGCCATACACATTTGTAGAGGTTTTACTTGCTTTAAAAAAC  
CTCCACACCTCCCCCTGAACCTGAAACATAAAATGAATGCAATTGTTGTTGT  
TAACTTGTTTATTGCAGCTTATAATGGTTACAAATAAAGCAATAGCATCACAA  
ATTCACAAATAAAGCATTTTTTTCACTGCATTCTAGTTGTGGTTTGTCCAAAC  
TCATCAATGTATCTTAA

TCATCAATGTATCTTAA

[illegible]

SEQ ID NO:2 SV40Xba#2 AAAATCTAGATCATAATCAGCCATACC

SEQ ID NO:3 SV40SacII **CCGCGGTTAAGATACATTGATGAGTTTGG**

SEQ ID NO:4 SV40Bam#2 AAAAGGATCCATTGATGAGTTTGGACAAACC

SEQ ID NO:5 IE1-3.5'      **ACTAGTTATCTCCATGATGGGCGCG**

SEQ ID NO:6 IE1-6.5'      **AGATCTATATAGTTGCTGATGGGCGCG**

**Dm.para primers**

SEQ ID NO:8 Dm.para5' 5'- TGAGGAAGAACGCAGTTTGTTC -3'

SEQ ID NO:9 Dm.para3' 5'- CGGGCATAAAATGAAACCTCG -3'

### Dm.white primers

SEQ ID NO:10      White5'      5'- CGCTGTGACACATACTTTCTG -3'

SEQ ID NO:11      White3'      5'- GTCTTAGAGCCAGATATGCG -3'

### Dm.vha55 (subunit B) primers for 3'UTR

SEQ ID NO:12      DmvB5'      5'- TCTCCACCTCCTGCAATATCCG -3'

SEQ ID NO:13      DmvB3'      5'- CCCATTCACTCTTGTGACCAGAG -3'

### Dm.vha26 (subunit E) primers for 3'UTR

SEQ ID NO:14      DmvE5'      5'- ACCAGAAAGAGAACCAGCATCAAC -3'

SEQ ID NO:15      DmvE3'      5'- ACCTGCCAGCGGTCTGTAAAAG -3'



1. **Introduction**

SEQ ID NO:16 para5' 5'- TGCATGGAATTGGCTTGACTTC -3'

SEQ ID NO:16 para5' 5'- TGCATGGAATTGGCTTGACTTC -3'

SEQ ID NO:17      para3'      5'- AGCACCAGTTGATAGAGATTCTCCC -3'

SEQ ID NO:18      Ms.hb.5'      5'- CTCGTTCTTATTCCTCCTAAC -3'

SEQ ID NO:18      Ms.hb.5'      5'- CTCGTTCTTATTCCTCCTAAC -3'

SEQ ID NO:19      Ms.hb.3'      5'- ATGAACGGGTCGTTGTACC -3'

SEQ ID NO:20 fas5' 5'- TACTGCACCAGAAATGGAAGAGC -3'

SEQ ID NO:20 fas5' 5'- TACTGCACCAGAAATGGAAGAGC -3'

SEQ ID NO:21      *fas3'*      5'- ACGGGTTGGTTGTTCATAGCC -3'

SEQ ID NO:22 Ms.EcR.5' 5'- CGTGCAACGTGCTCGTTTTTAC -3'

SEQ ID NO:22 Ms.EcR.5' 5'- CGTGCAACGTGCTCGTTTTTAC -3'

SEQ ID NO:23      Ms.EcR3'      5'- TTAGGAGTTGTAGGAGGCATCGG -3'

SEQ ID NO:24      MsvB5'      5'- GATCTGGTTTCGATTGTTTCCG -3'

SEQ ID NO:24      MsvB5'      5'- GATCTGGTTTCGATTGTTTCCG -3'

SEQ ID NO:25      MsvB3'      5'- CGAGGACCAACTCAATTTGGAATG -3'

SEQ ID NO:26      MsvE5'      5'- GGTGACCCACATTCACCTCGTTATAC -3'

SEQ ID NO:26      MsvE5'      5'- GGTGACCCACATTCACCTCGTTATAC -3'

SEQ ID NO:27      MsvE3'      5'- AACCTACAGACCTCAATGCCTCC -3'

## Figure 9

**Portion of Multiple Cloning Site from Plasmid PCR2.1 Incorporated into Modified Baculovirus Transfer Plasmids**

SEQ ID NO:7

AGCCGAATTCTGCAGATATCCATCACACTGGCGGCCGCTCGAGCATGCATCT  
AGA

09220463-081304  
T0CT80-09482660